



OIPE

RAW SEQUENCE LISTING

FATENT AFFLICATION: US/09/835,072

DATE: 17 - 10 + 3 TIME: 13:00:10

Input Set : A:\IS4584.txt

Output Set: N:\CRF4\02032003\1835072.raw

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4 <110 - APPLICANT: Goodlett, David R.
        Keller, Andrew
 7 K120. TITLE OF INVENTION: METHODS FOR QUANTIFICATION AND DE NOVO
        FOLYPERTIDE SEQUENCING BY MASS SPECTROMETRY
11 01300 FILE REFERENCE: PHIS 4584
15 <1400 CURRENT APPLICATION NUMBER: US 09/835,072
14 <141 - OUFFENT FILING DATE: 2001-04-13
16 41600 NUMBER OF SEQ ID NOS: 14
1: <17 < SOFTWARE: FastSEQ for Windows Version 4.0
20 <2100 (550 ID NO: 1
31 42110 LENGTH: 10
21 H21HH TYFE: PHT
23 C218 DESANISM: Homo sapiens
SE 440 % SEQUENCE: 1
16 3lm Asp Val Asp Asm Ala Ser Leu Ala Arg
17 1
                                        10
51 K21 0 SEP ID NO: 2
31 42111 LENGTH: 7
32 K21LD TYPE: PRI
33 40114 OFGANISM: Homo sapiens
35 K40 CH SEQUENCE: 2
Ge Gin Asp Val Asp Ash Ala Ser
37 1
40 00100 SEQ ID NO: 3
41 -(211) LENGTH: 9
40 -00100 TYPE: PRT
43 -021%: CFGANISM: Homo sapiens
45 HAORD SEQUENCE: 3
46 Bin Gln Tyr Glu Ser Val Ala Ala Lys
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50 00100 SEQ ID NO: 4
51 <2211 : IENGTH: 13
52 K2110 TYPE: PRT
53 <2130 CRGANISM: Homo sapiens
55 K4000 SEQUENCE: 4
56 Gln Glu Tyr Asp Glu Ser Gly Pro Ser Ile Val His Arg
                                        10
60 kalom seg id Mo:
61 <211 - LENGTH: 8
62 FAIL TYPE: PPT
6 + 213 + ORGANICM: Hor - saylens
CE +400 - SEQUENCE: E
ee Gin Glu Tyr Asy Glu Ser Gly Er
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INTEL CALL THE

FATENT AFFLI MATION: US/09/835,072 TIME: 13:06:10 Input Set : A:\IS4584.txt Curput Set: N:\CRF4\02032003\1835072.raw ni kalah seg in No: 6 ol (211) EENSTH: 1 72 - 212'- TYPE: PRT 73 K213 - ORGANISM: Homo sapiens 75 <400 - SEQUENCE: 6 $76~\mathrm{Ala}~\mathrm{Hy}~\mathrm{Phe}~\mathrm{Ala}~\mathrm{Gly}~\mathrm{Asp}~\mathrm{Asp}~\mathrm{Ala}~\mathrm{Fro}~\mathrm{Arg}$ 7. 1 80 3210 - SEQ ID NO: 81 <211 - LENGTH: 16 8. K212 - TYEE: PRT 83 KOls + ORGANISM: Homo sapiens 85 King - SEQUENCE: 7 86 Ser Tyr Glu Let Pro Asp Gly Gln Val Ile Thr Ile Gly Ash Glu Arg 10 90 Hally + SEQ ID NO: 8 91 WHILL LENGTH: 9 91 HOLL FIYES: PRT 9: 4313 - ORGANISM: Homo sapiens 91 -: 1000 - SEQUENCE: 8 96 Pro Asp Ash Ala Val Ile Thr Ile Gly 97 1 5 100 d. 11 - SEQ ID NO: 9 101 -1.11 - DENGTH: 11 102 0.10 0 TYPE: PRT 103 4.13 + OFGANISM: Homo sapiens 165 -0467 - SEQUENCE: 9 106 Leu Leu Leu Leu Gly Ala Gly Glu Ser Gly Lys 10 1 10 110 -000 SEQ ID NO: 10 111 -Mill LENGTH: 8 100 HARLE TYPE: PRT 113 - 219 - OFGANISM: Homo sapiens 115 -: 40: - SEQUENCE: 10 116 Leu Leu Leu Gly Ala Gly Glu 11 % -00100- SEQ ID NO: 11 1:1 -:::::: LENGTH: 12 11.0 + .11.0 TYPE: PFT 115 - 2150 OFGANISM: Homo sapiens 11 to 4000 SEQUENCE: 11 110 Gly Ash Leu Gir Ile Asp Phe Ala Asp Pro Ser Arg 130 -Chlor SEQ ID No: 12 131 KU111: LENGTH: 5 13. K2000 TYPE: PAT 133 <213> ORGANISM: Homo sapiens 13. <400> SEQUENCE: 12 1st Ile Asp Phe Ala Asp

RAW SEQUENCE LISTING

Input Set : A:\IS4584.txt

Ourput Set: N:\CRF4\02032003\I835072.raw

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147 1 5
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Impur Set : A:\IS4584.txt

Output Sat: N:\CRF4\02032003\1835072.raw